

Figure 1: Yeast Genome; One Mismatch; Truncation to length "L"  
Theory and Measurements

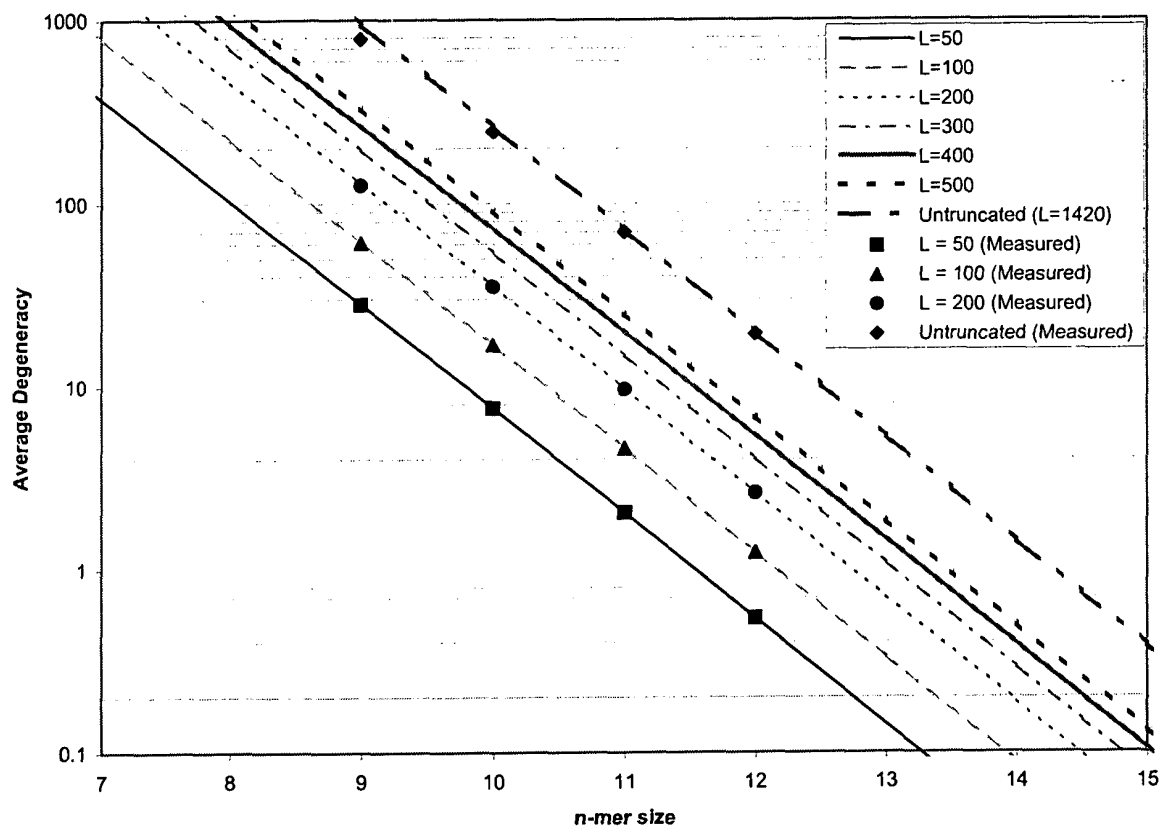


Figure 2: Mouse Genome; One Mismatch; Truncation to Length "L"

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Theory and Measurements

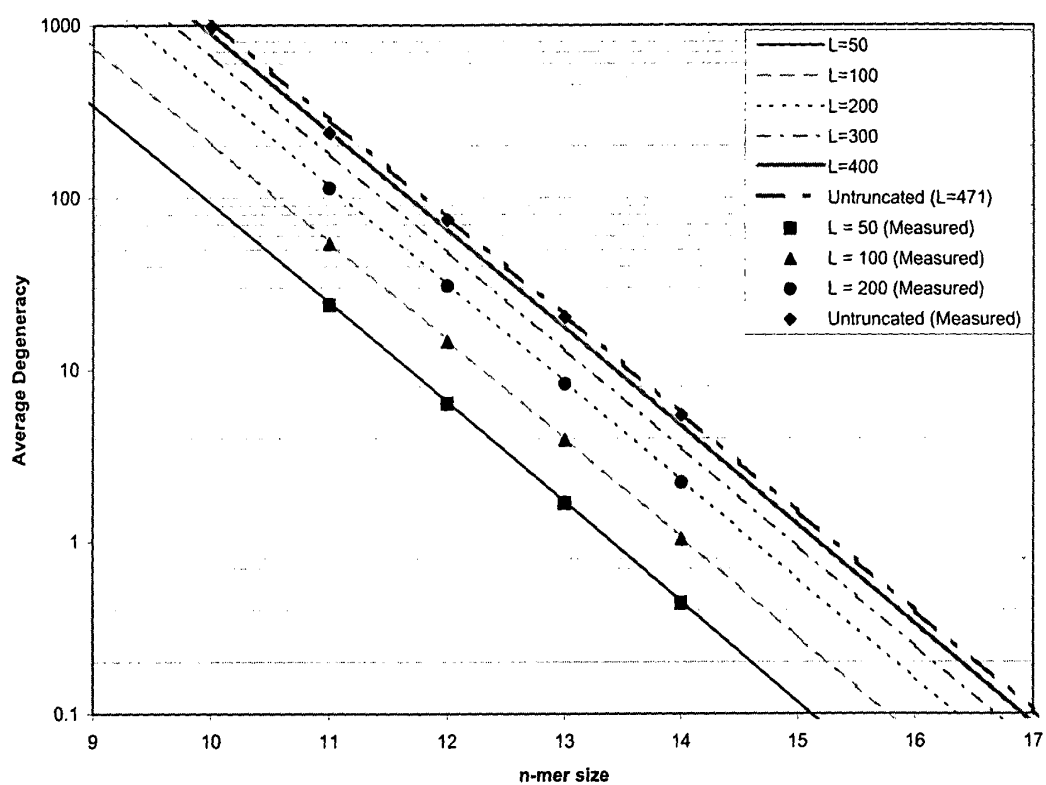


Figure 3: Relationships between parameter values such that average degeneracy = 1 (theoretical predictions)

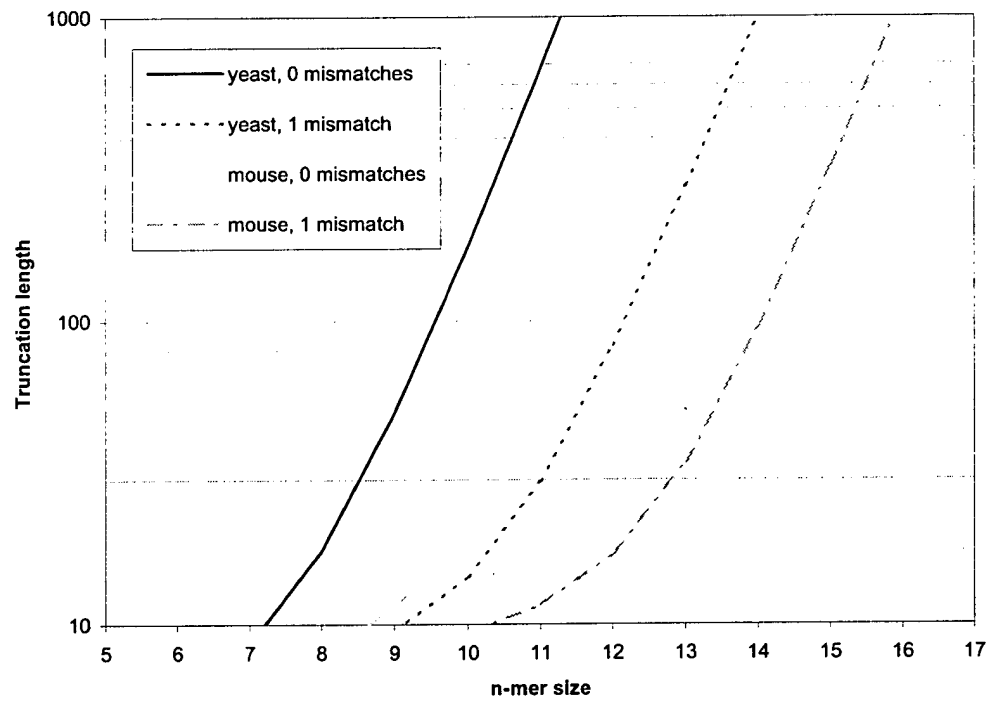


FIG. 4A

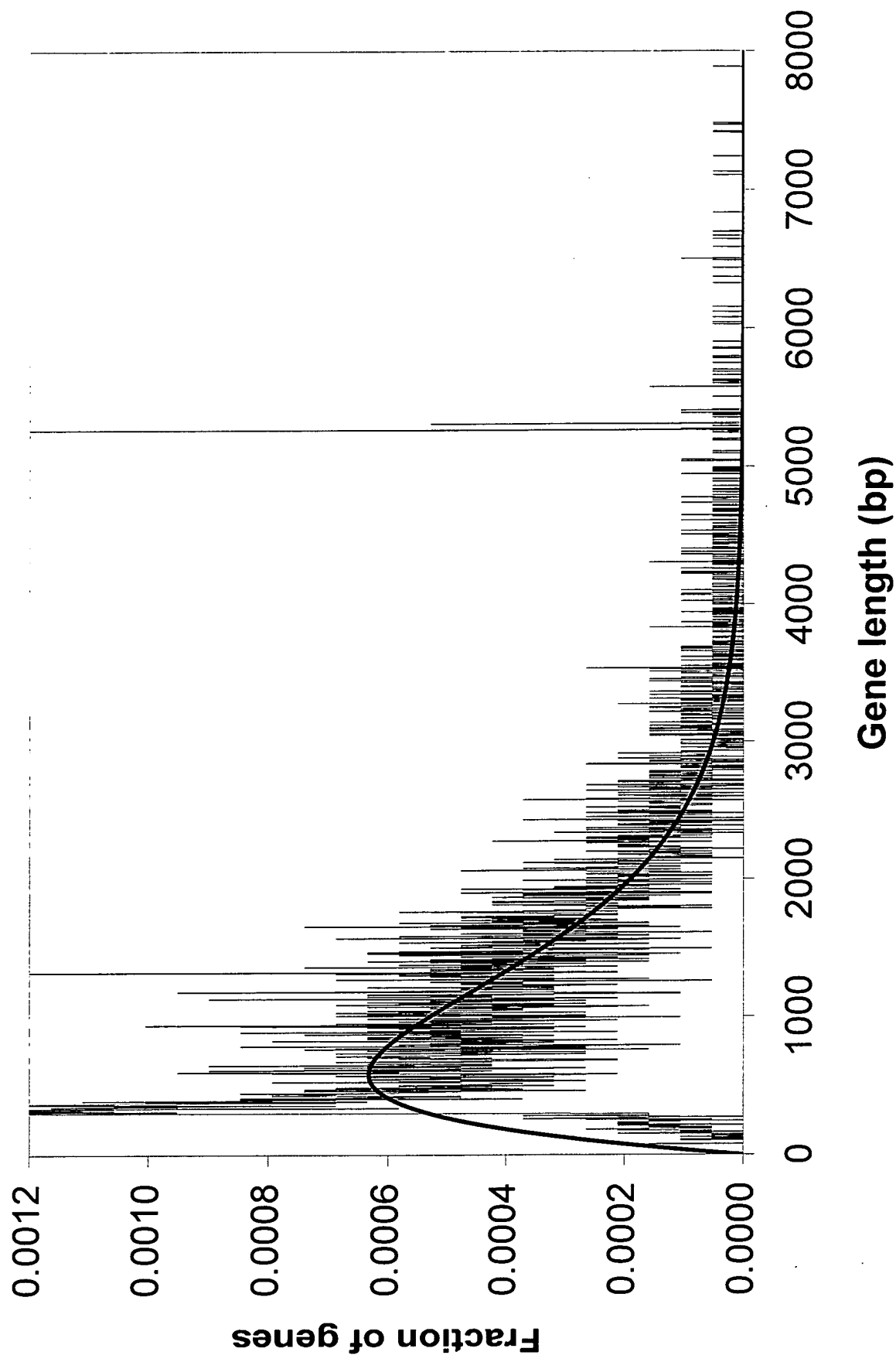


FIG. 4B

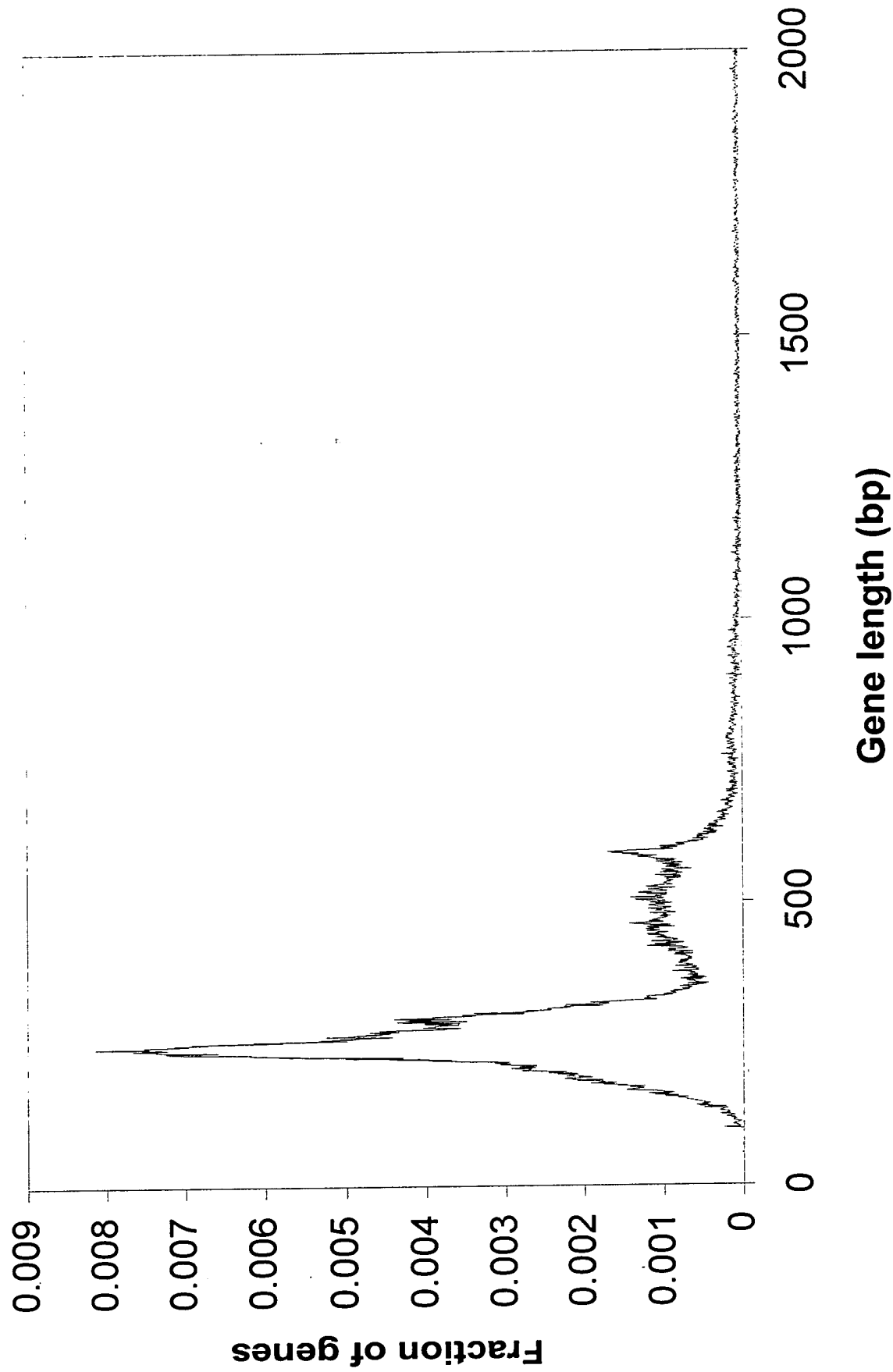


FIG. 5A

8-mers; mismatches: 0

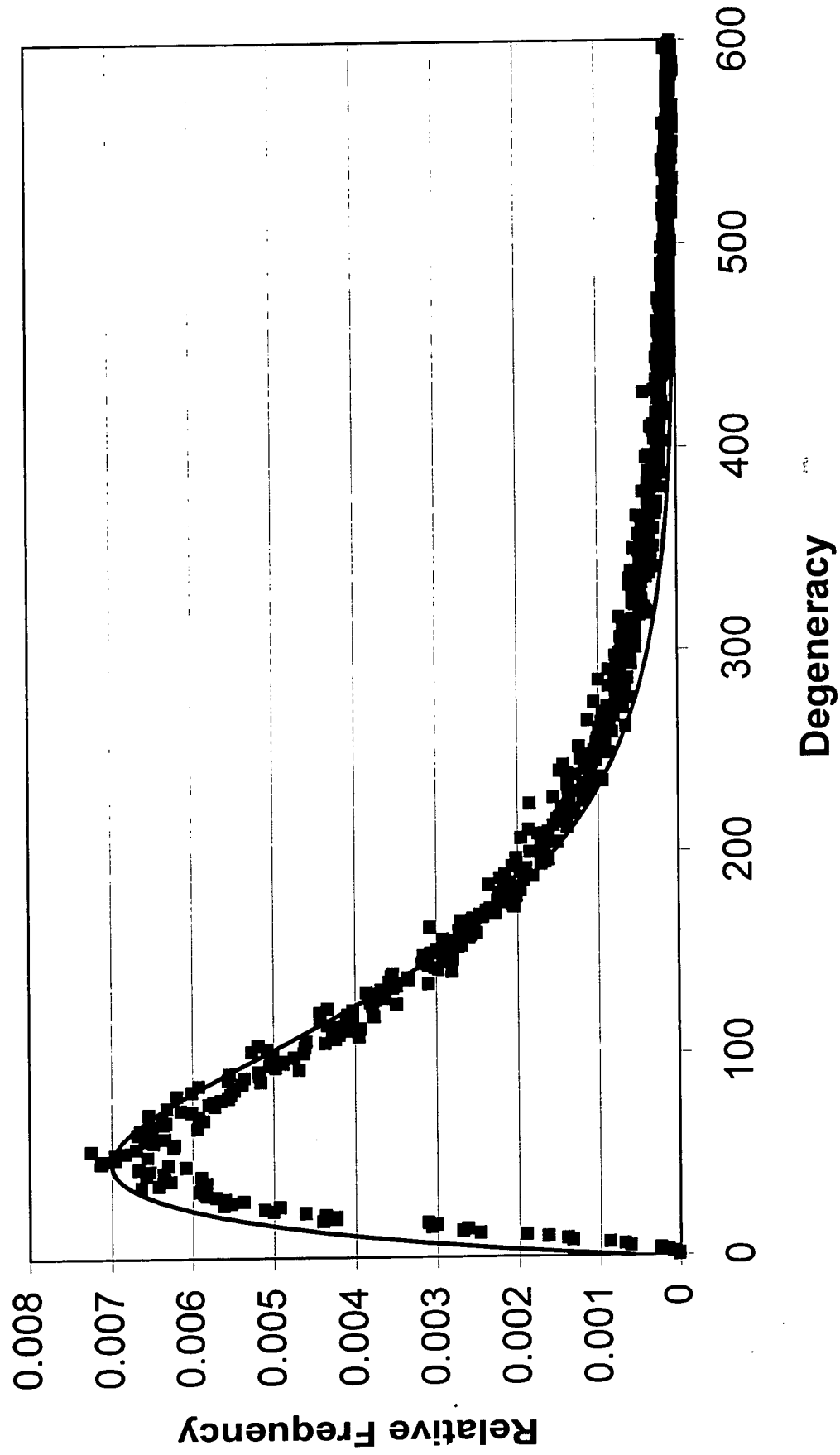


FIG 5B

8-mers; mismatches: 1

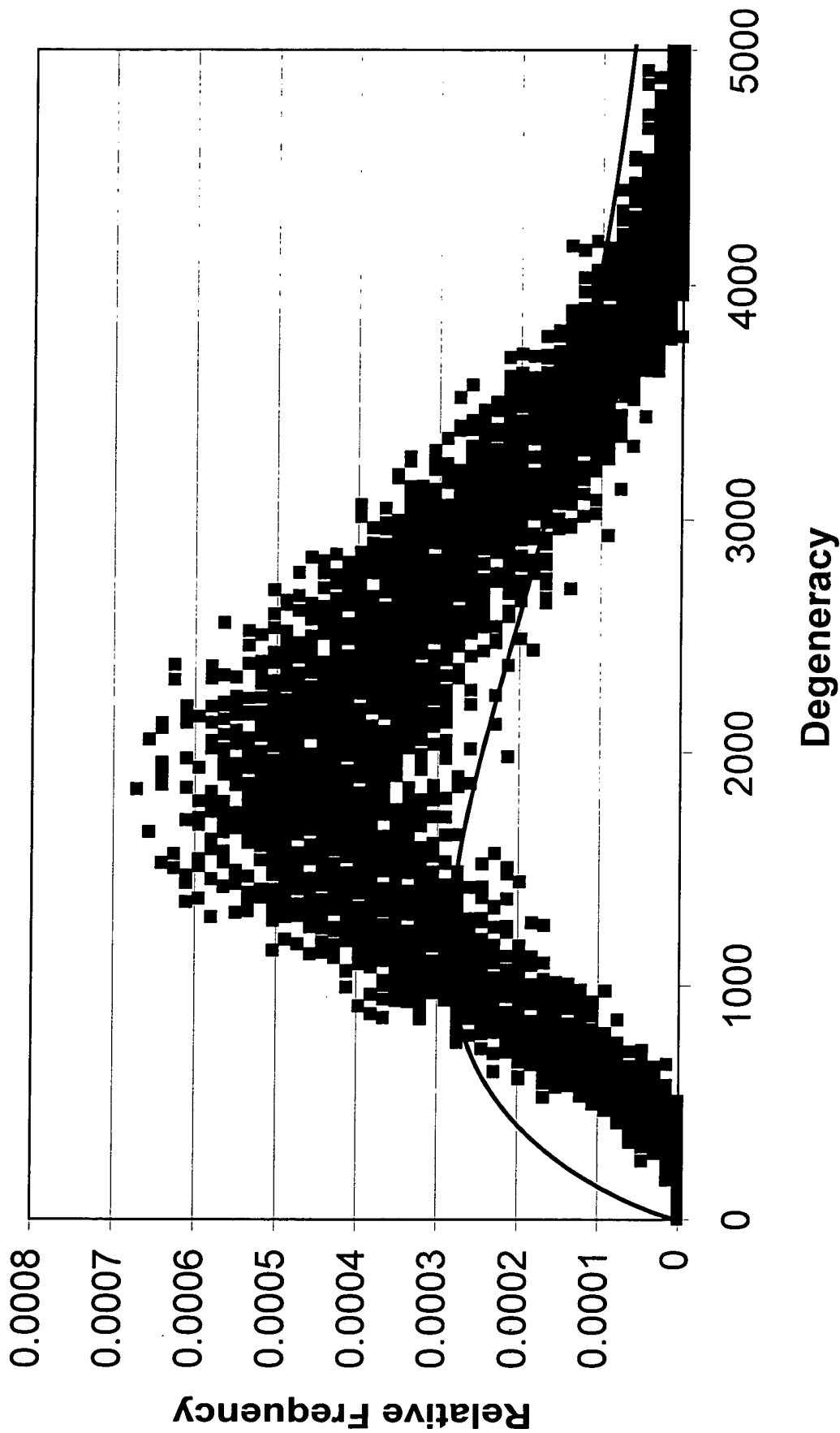


FIG. 5C

9-mers; mismatches: 0

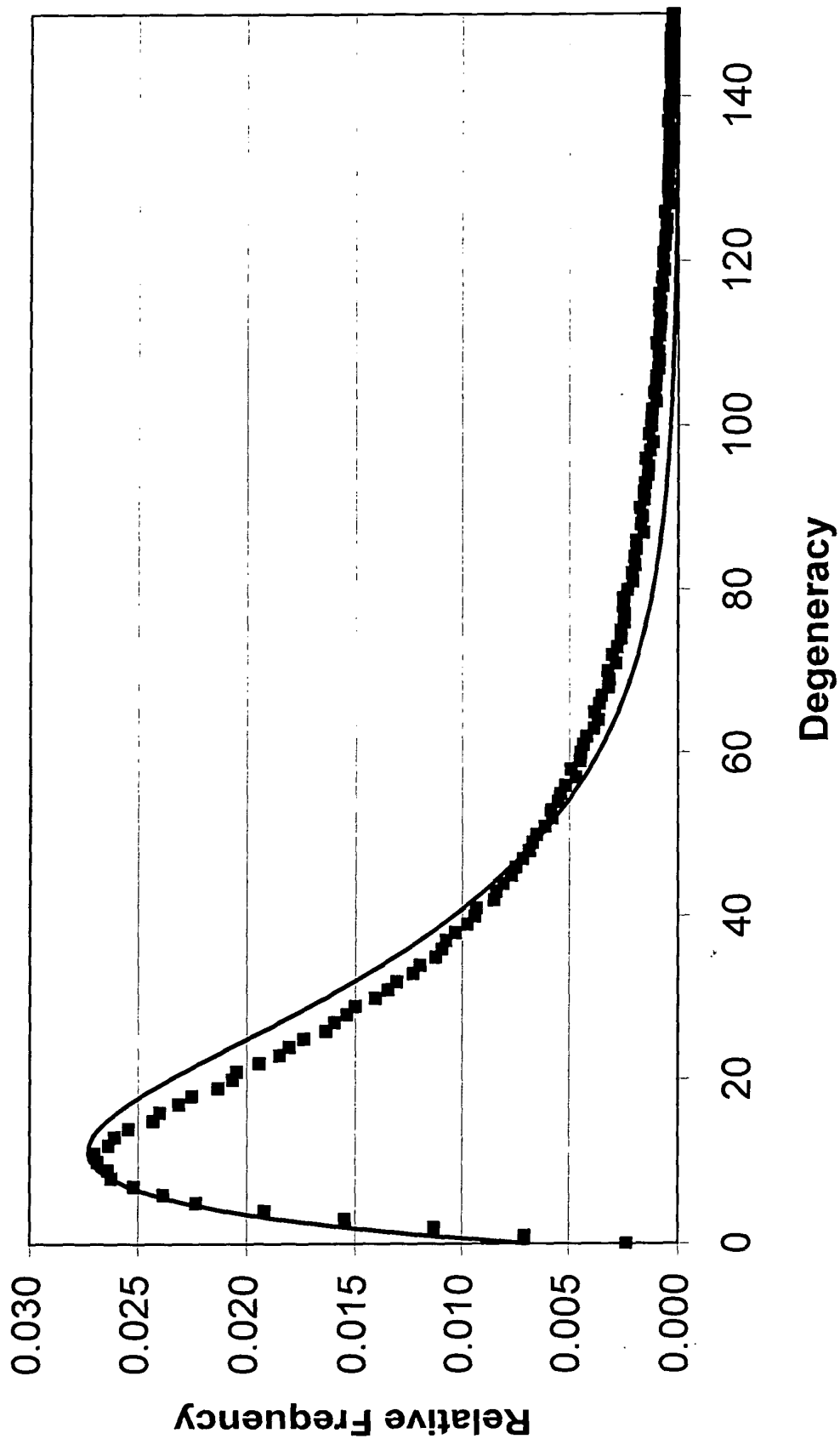




FIG. 5D

9-mers; mismatches: 1

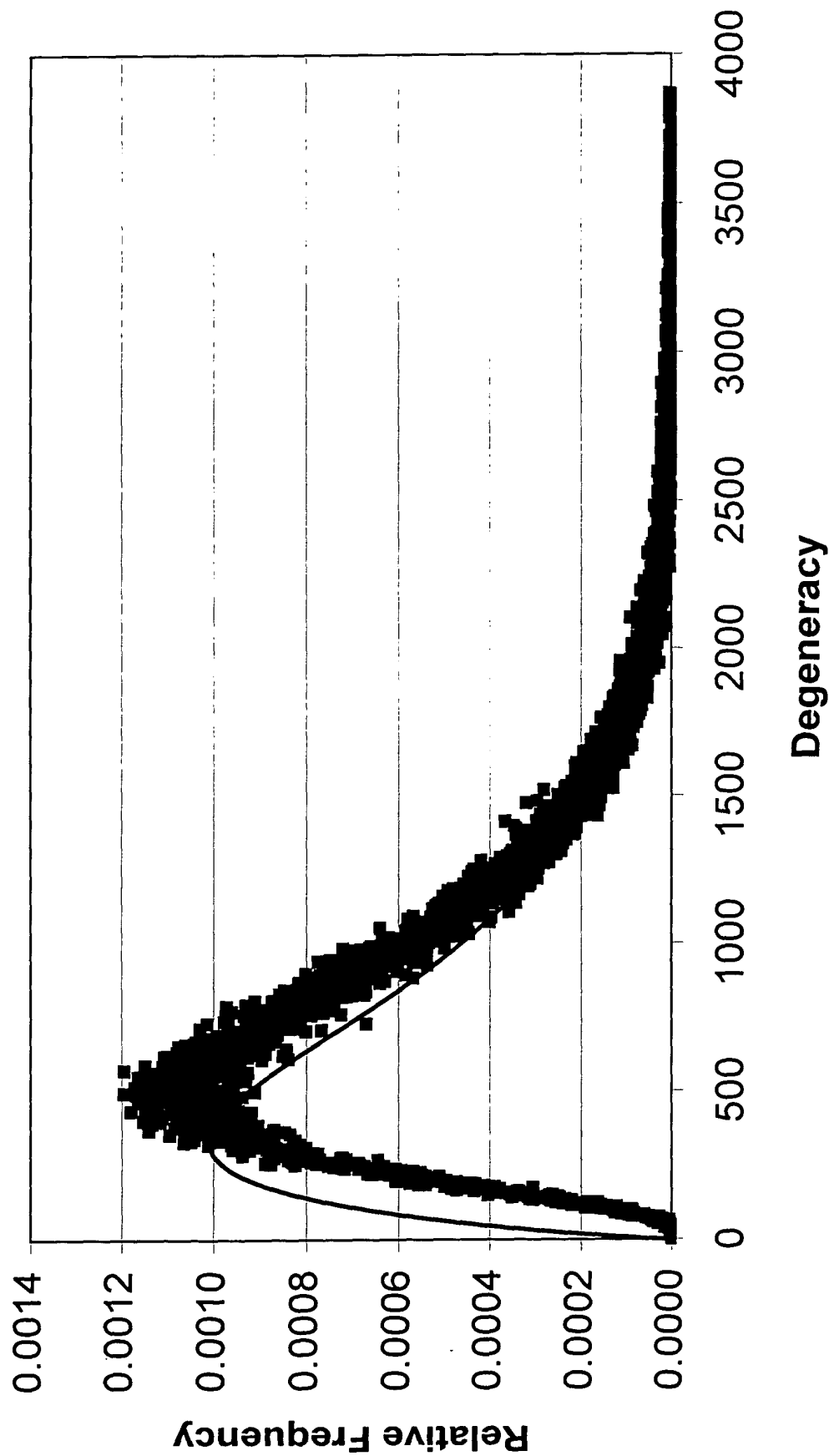
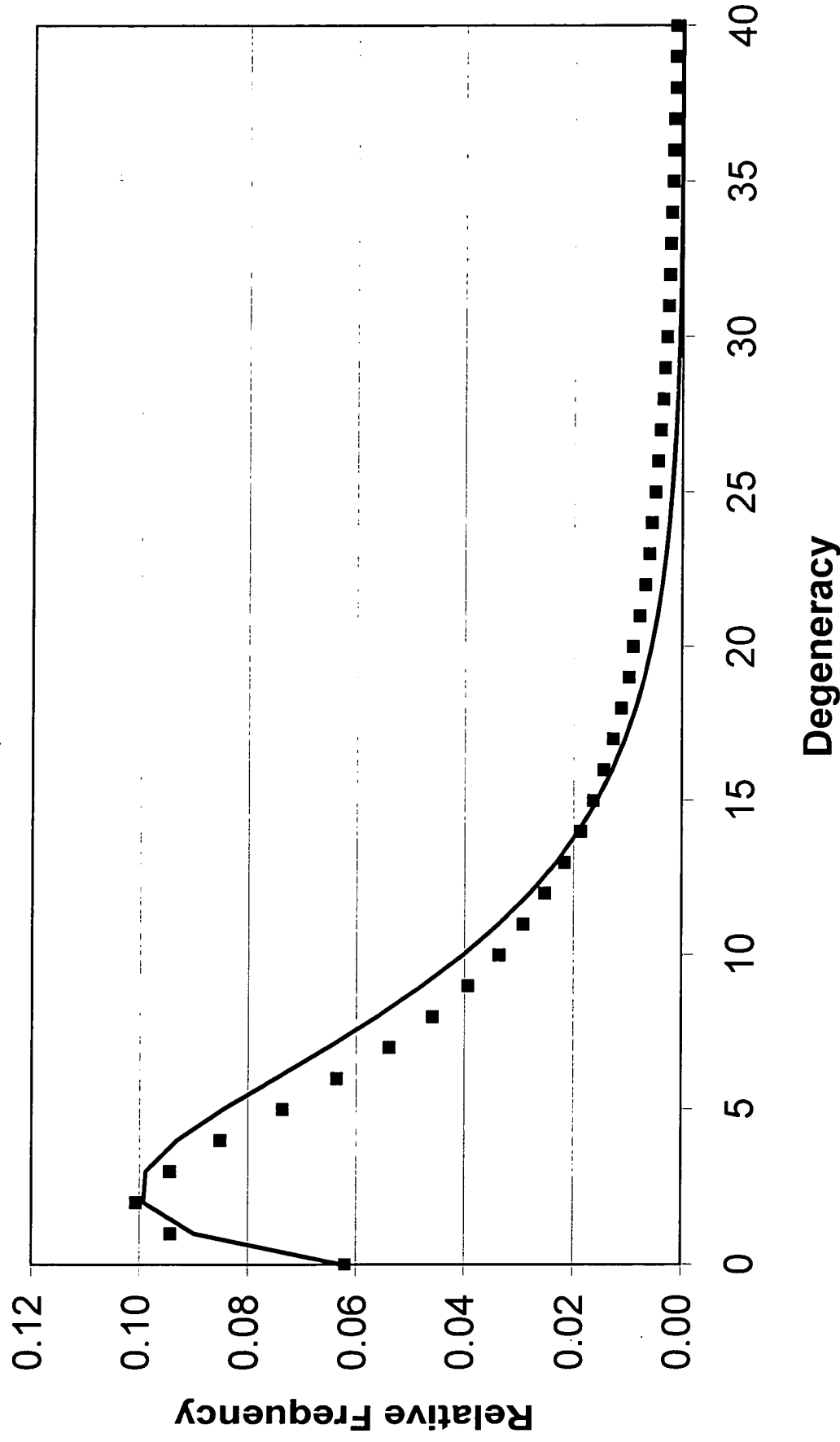


FIG 5E

10-mers; mismatches: 0



10-mers; mismatches: 1

FIG. 5F

10-mers; mismatches: 1

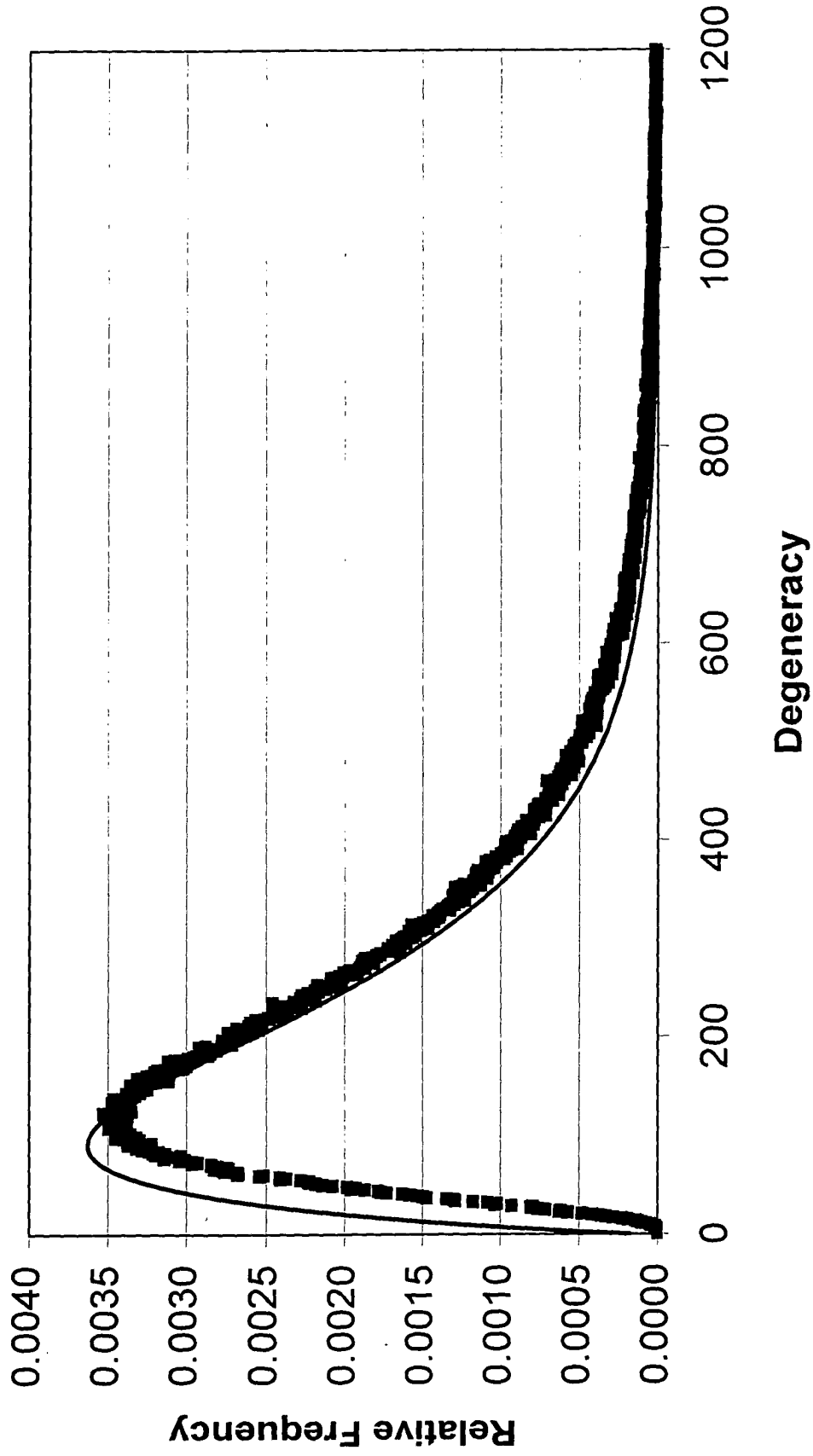


FIG. 5G

11-mers; mismatches: 0

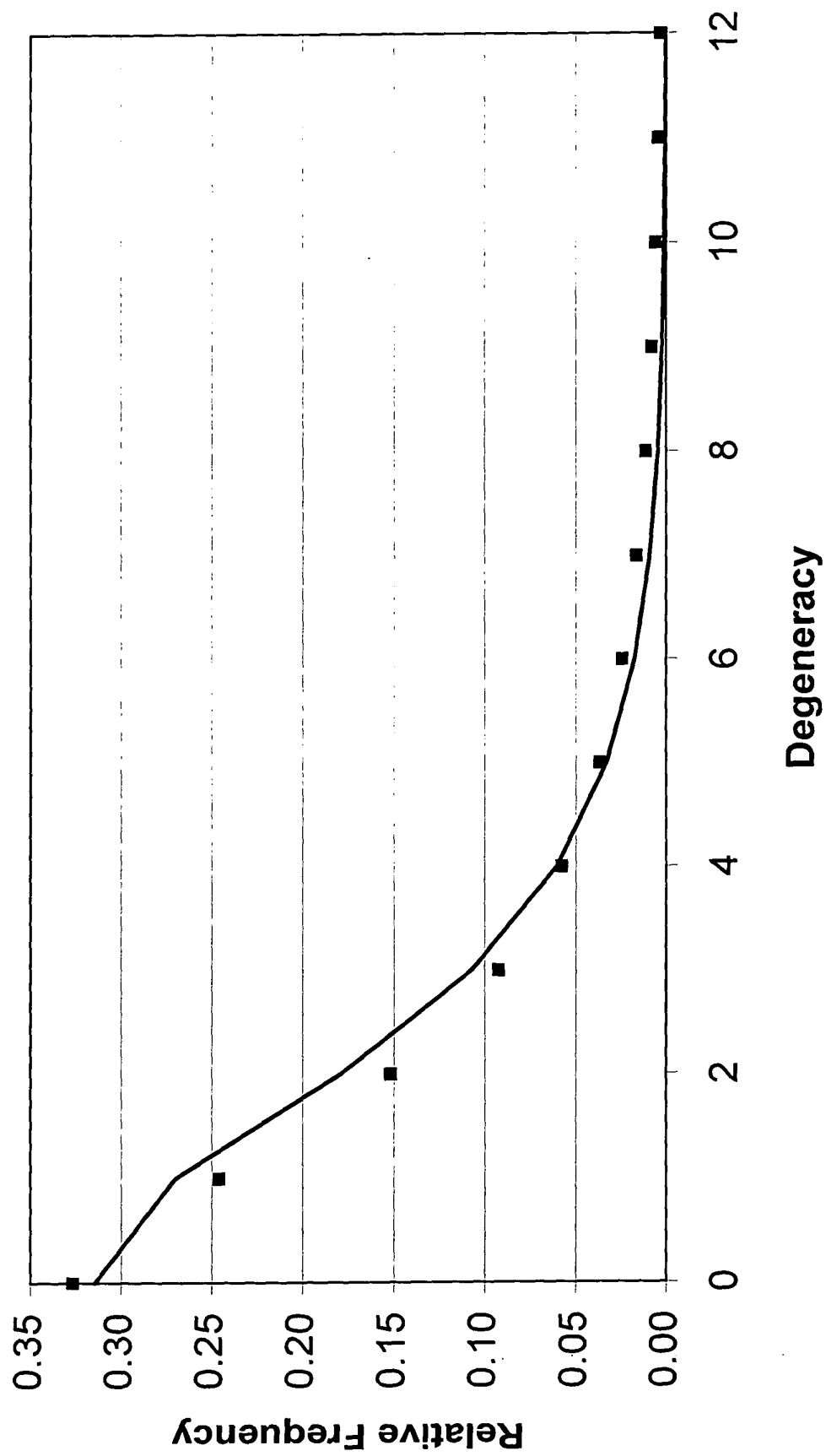
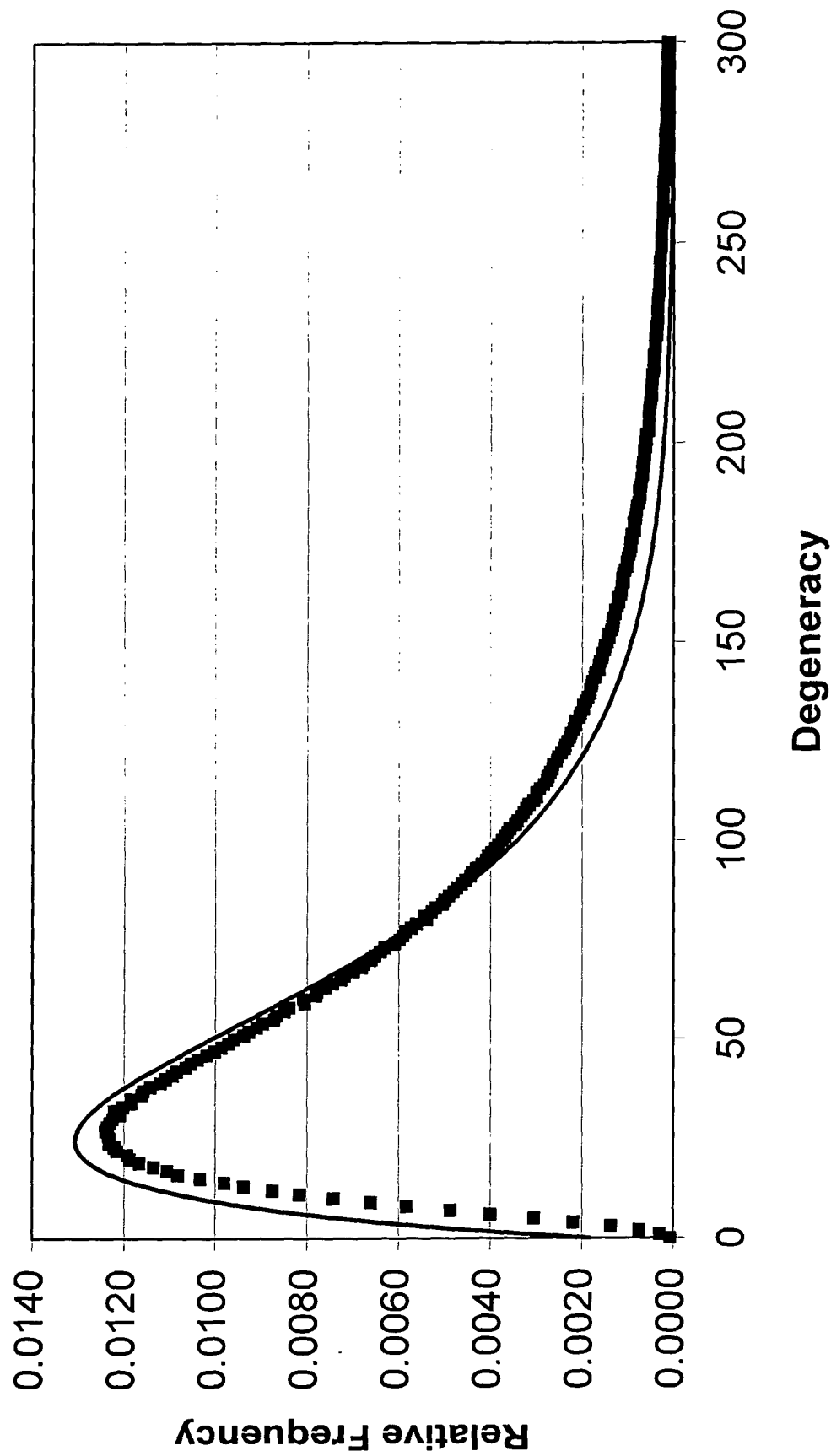


FIG. 5H

11-mers; mismatches: 1



12-mers; mismatches: 0

FIG. 5I

12-mers; mismatches: 0

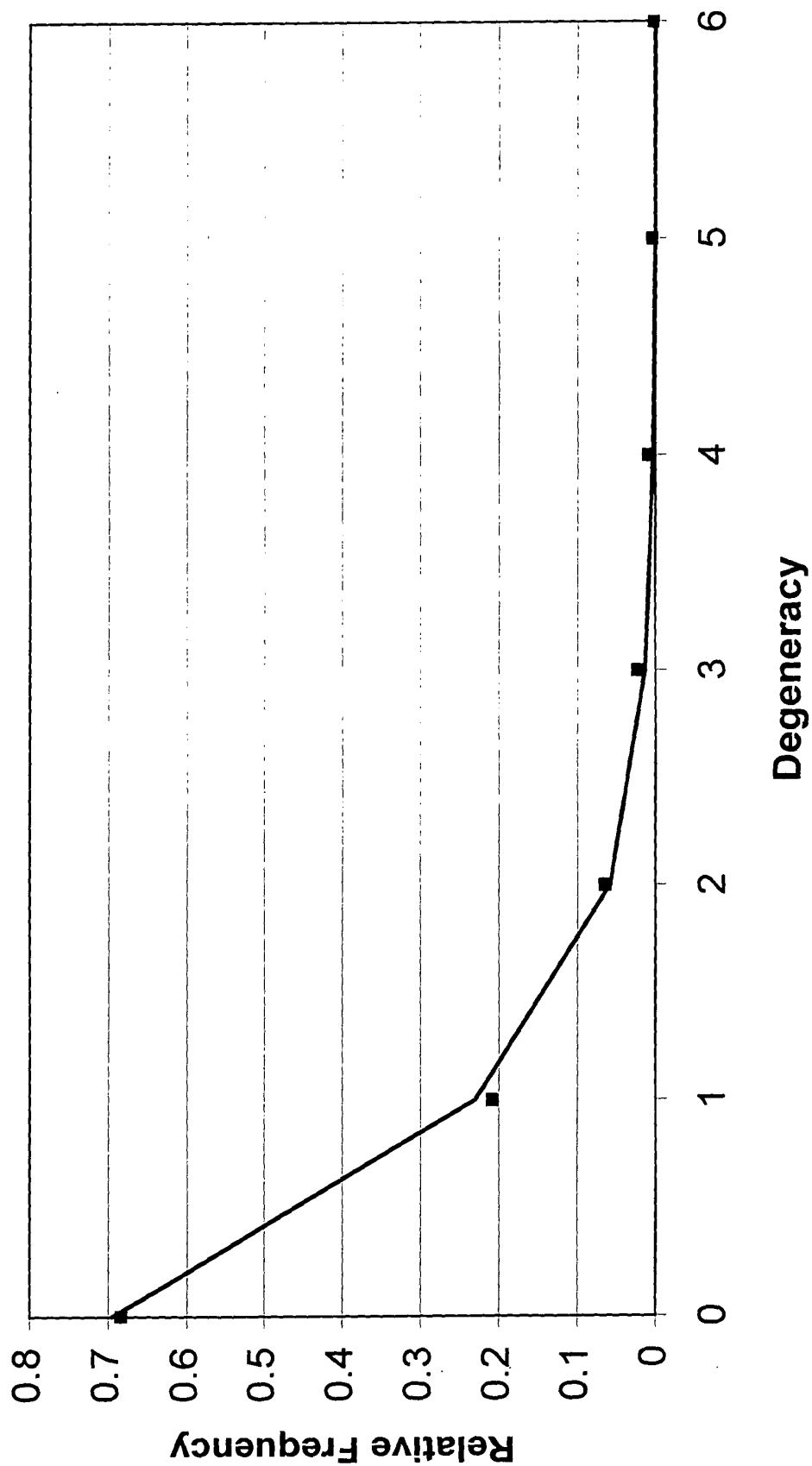


FIG. 5J

12-mers, mismatches: 1

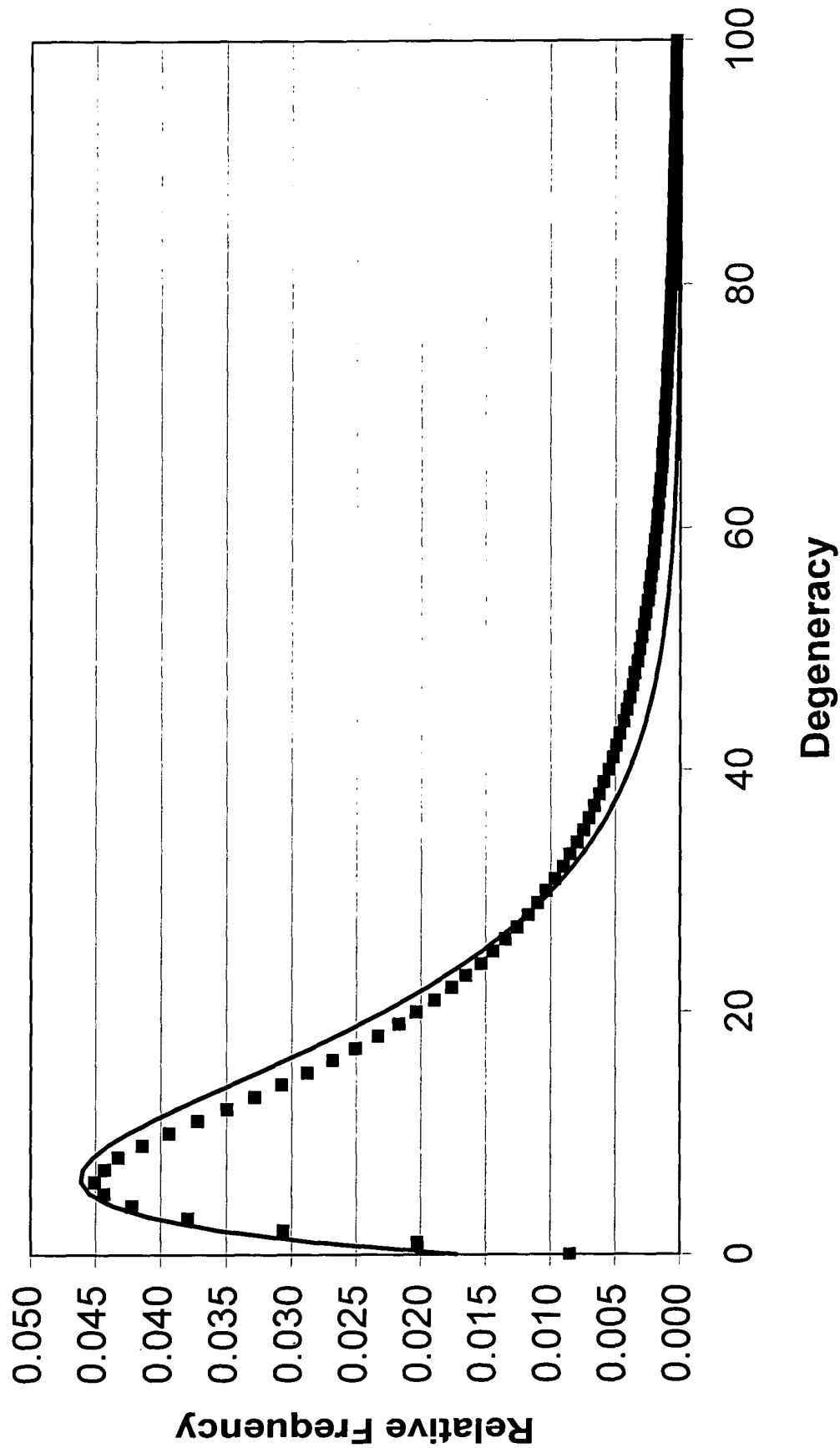


FIG. 6A

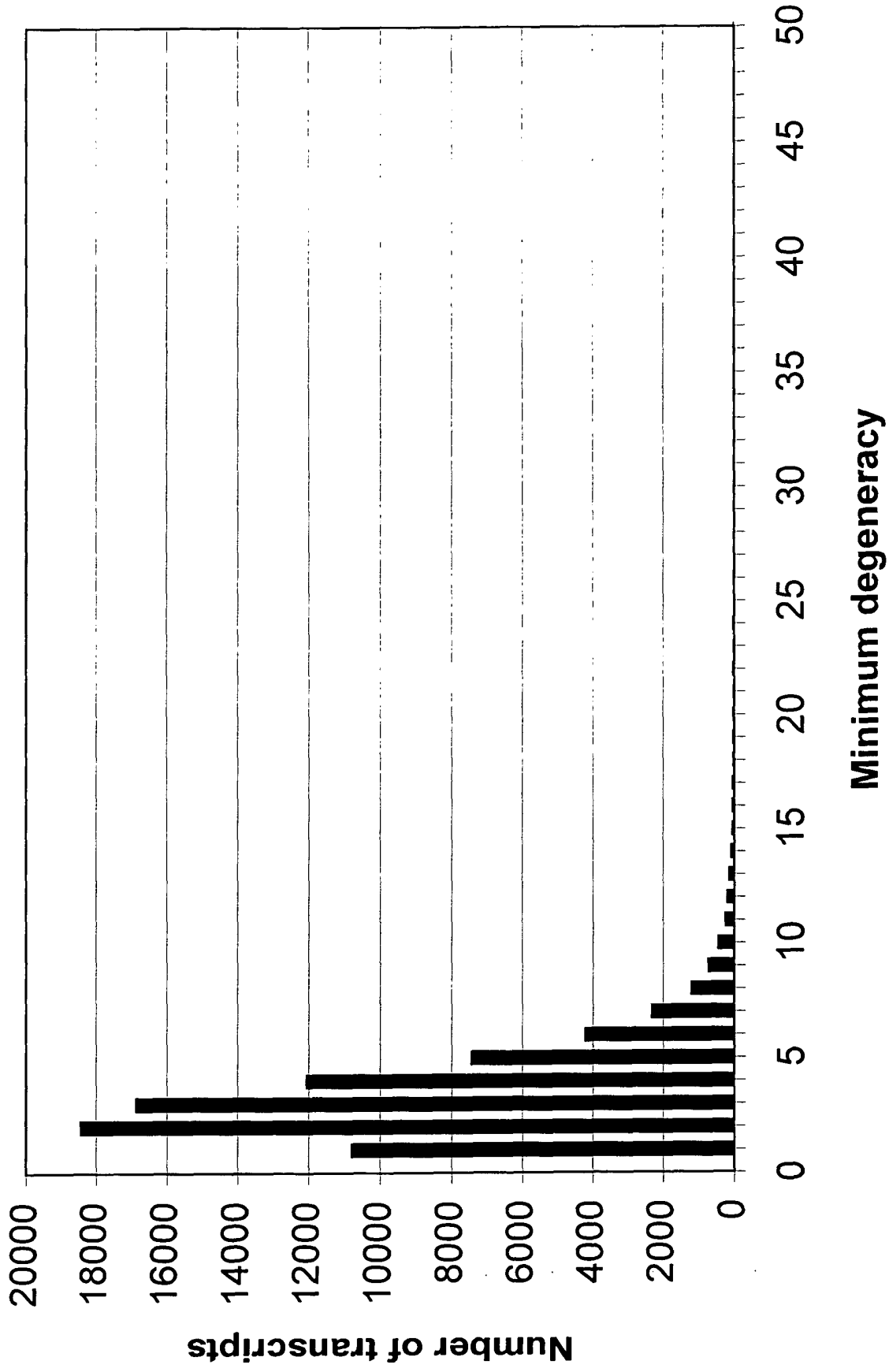




FIGURE 4B

FIG. 4B

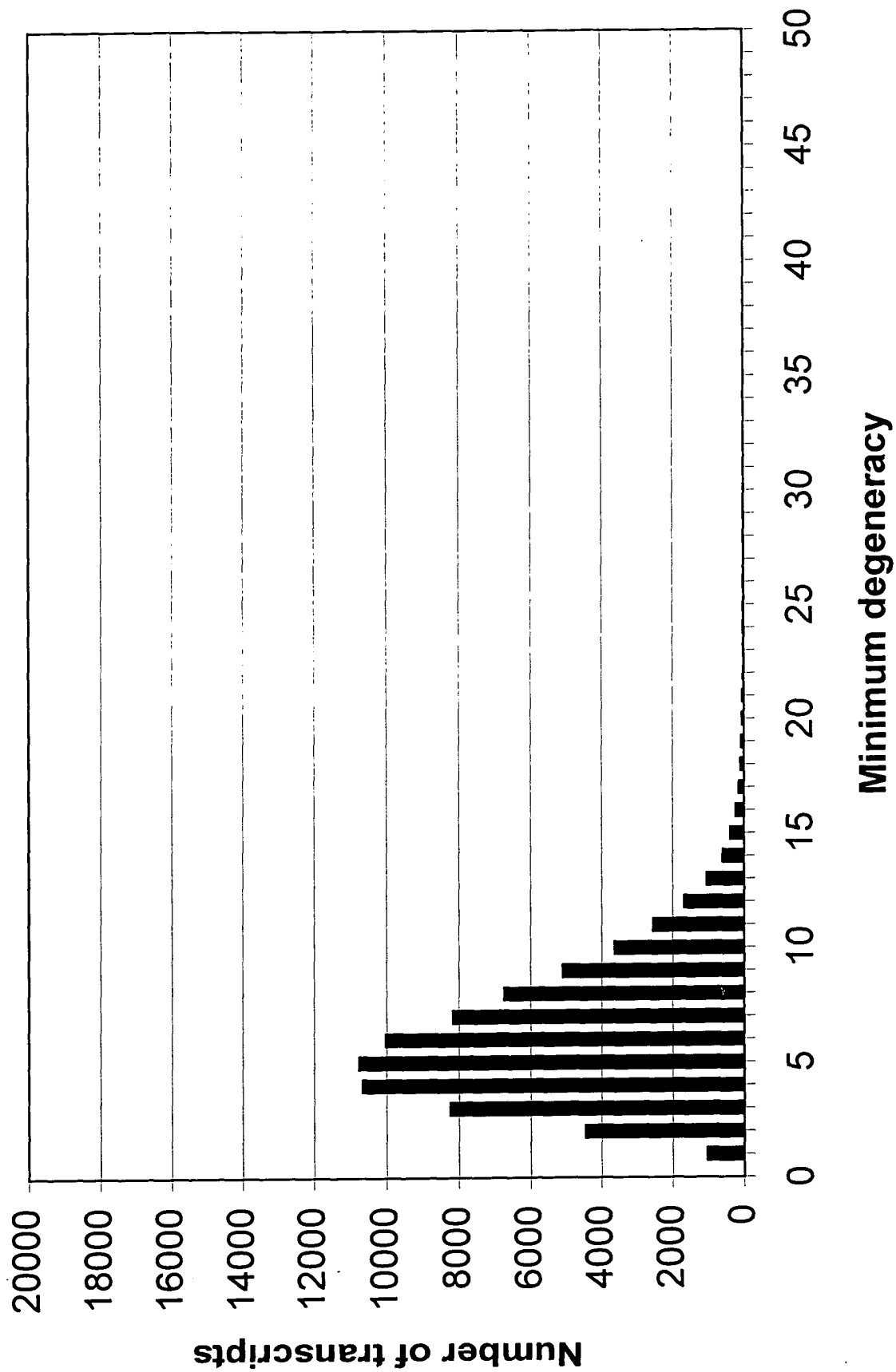


TABLE 5-100000

FIG. 6C

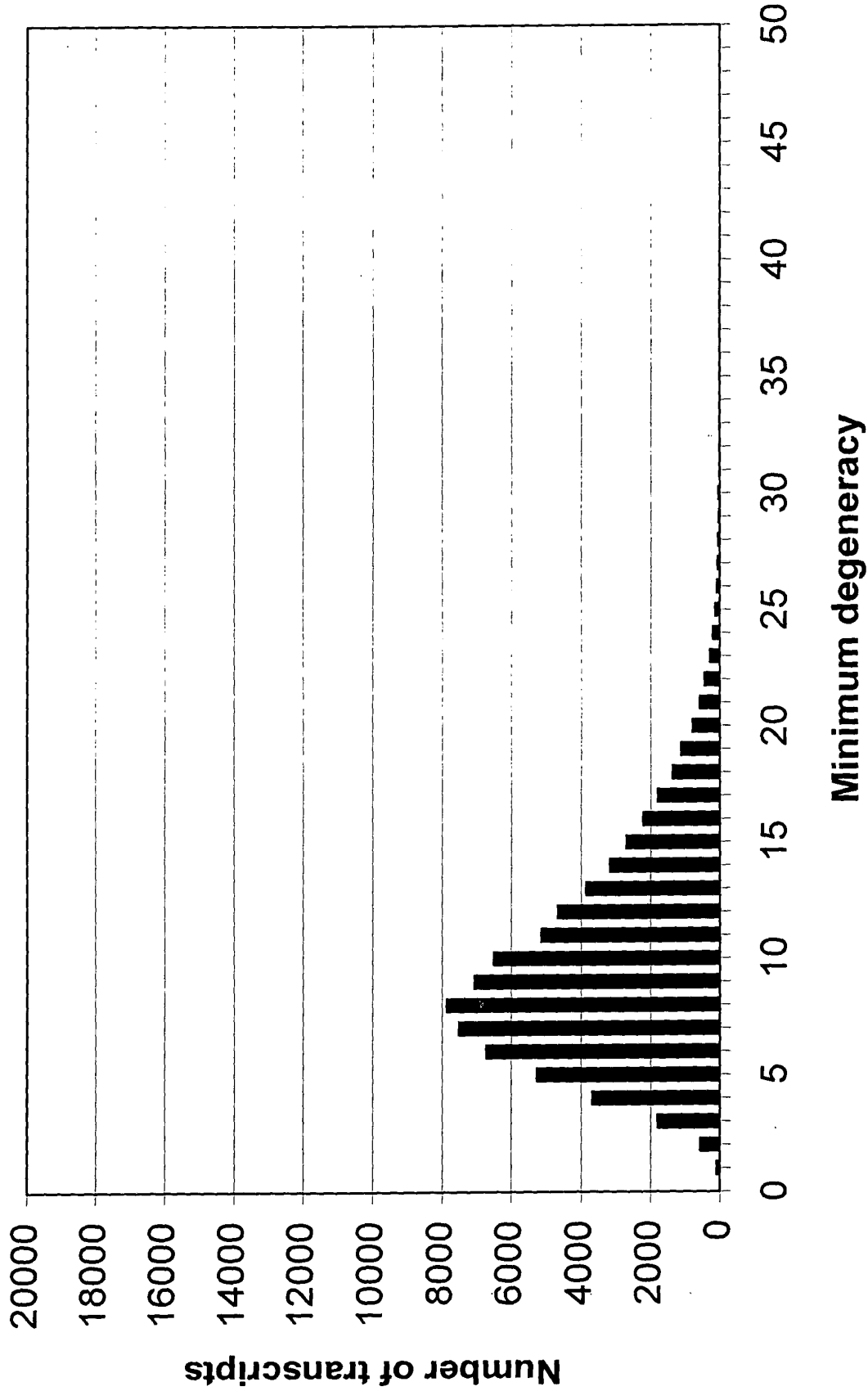


FIGURE 3

FIG. 4D

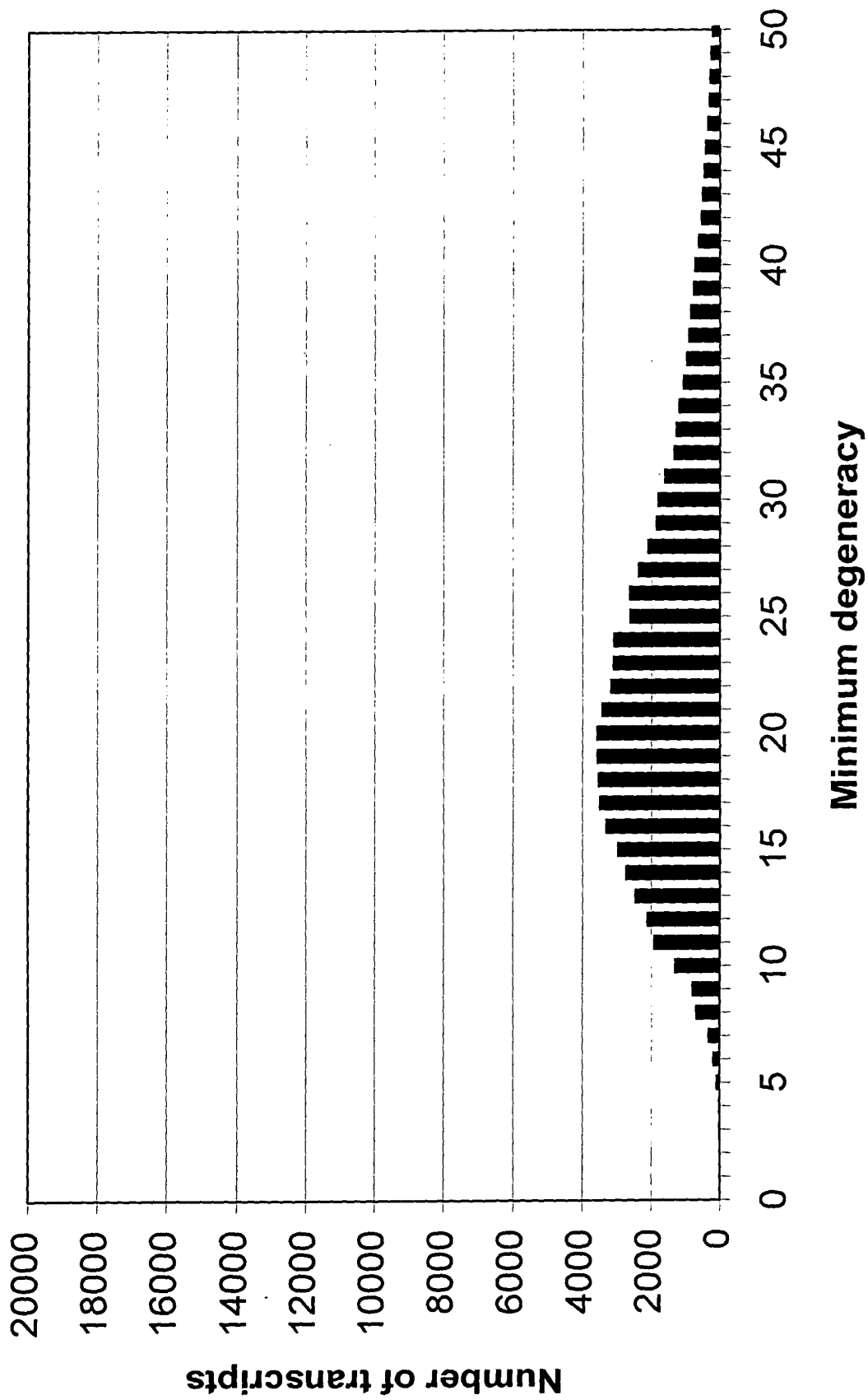
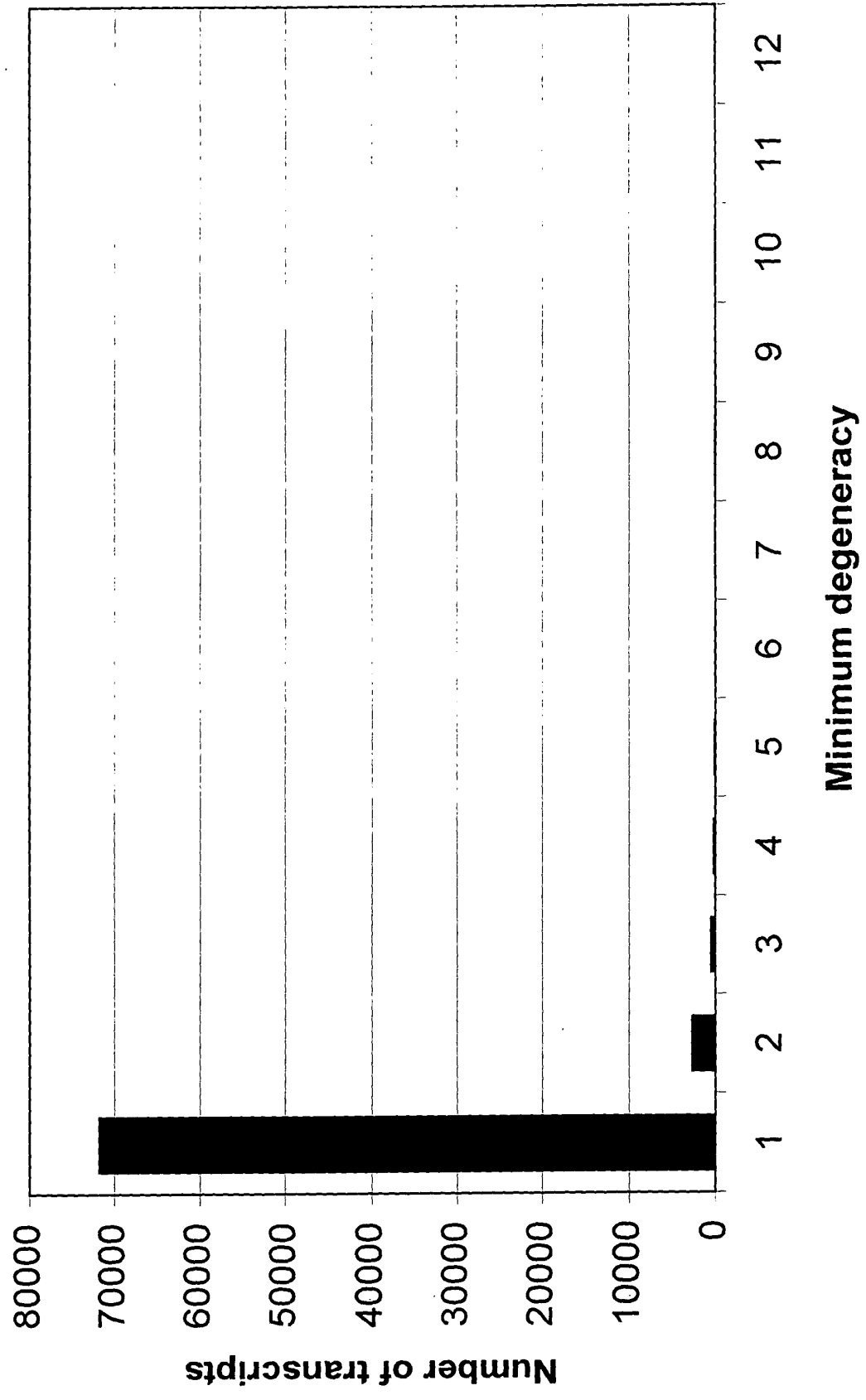


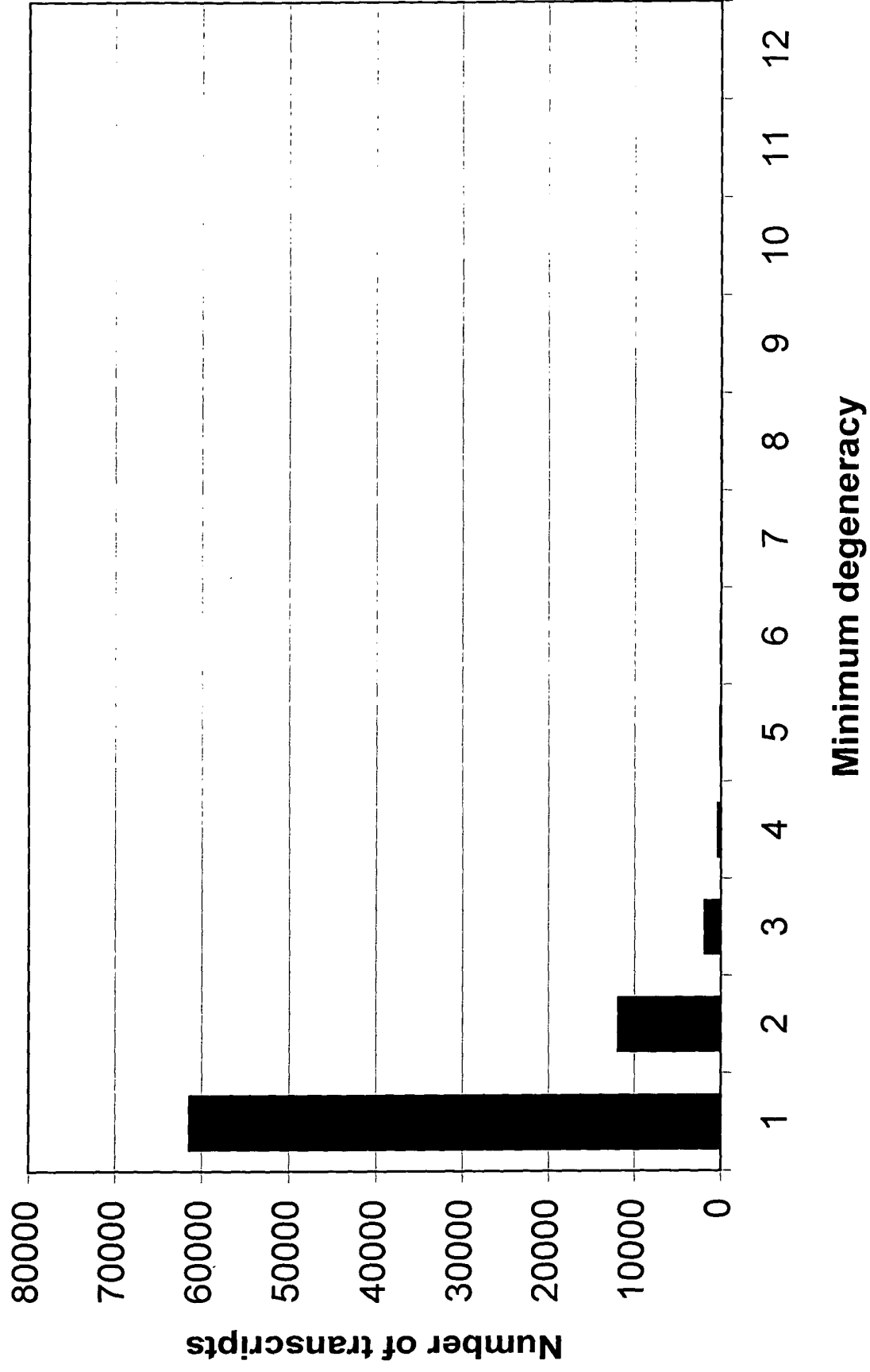
Figure 1

FIG. 1

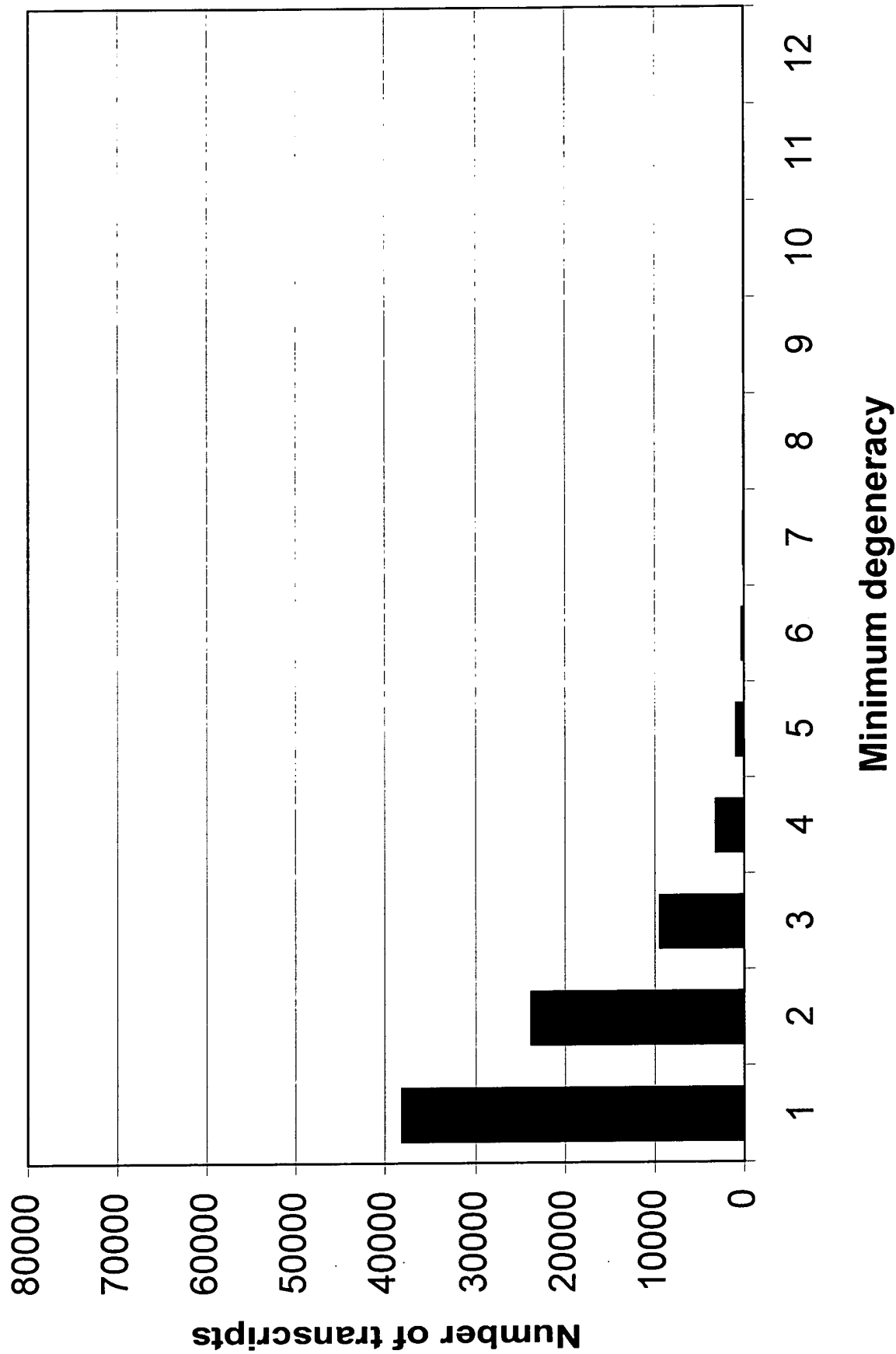


FOR RELEASE

FIG. 6F



total 340000  
FIG. 6G



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FIG. 6H

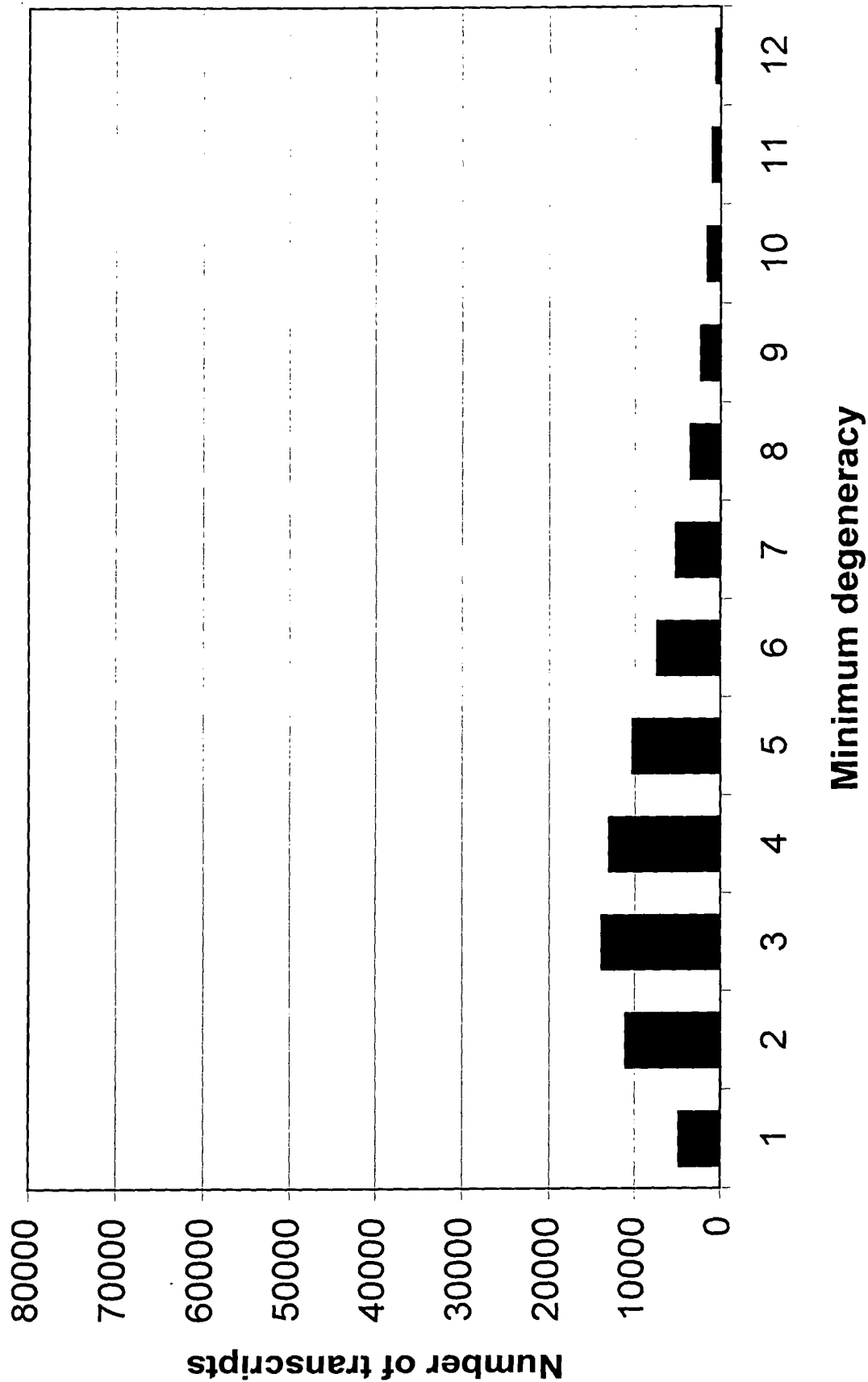


FIG. 7A

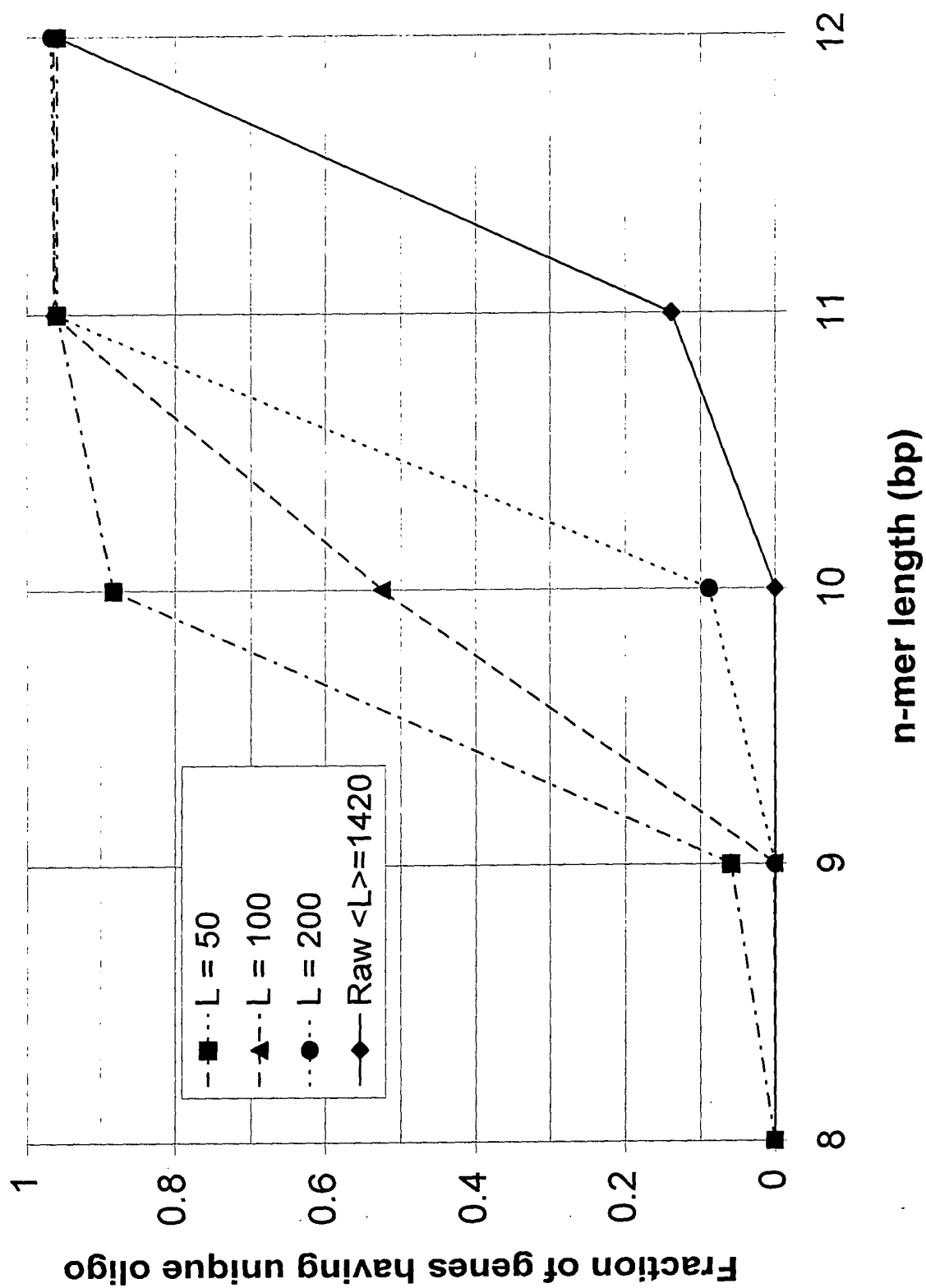




FIG. 7B

